

**AMENDMENTS TO THE CLAIMS**

Please enter the following amendments without prejudice or disclaimer.

This listing of claims will replace all prior versions, and listings, of claims in the application.

**In the claims:**

1-23. (Canceled)

24. (Currently amended) A method of reducing or moderating a postprandial rise in plasma glucose in a mammal comprising administering to said mammal an amylin or an amylin agonist in an amount effect effective to reduce or moderate a postprandial rise in plasma glucose, wherein the amylin agonist is a peptide.

25. (Previously presented) The method of claim 24 wherein the amylin agonist is an amylin agonist analogue having the following amino acid sequence:

<sup>1</sup>A<sub>1</sub>-X-Asn-Thr-<sup>5</sup>Ala-Thr-Y-Ala-Thr-<sup>10</sup>Gln-Arg-Leu-B<sub>1</sub>-Asn-<sup>15</sup>Phe-Leu-C<sub>1</sub>-D<sub>1</sub>-E<sub>1</sub>-<sup>20</sup>F<sub>1</sub>-G<sub>1</sub>-Asn-H<sub>1</sub>-G1y-<sup>25</sup>Pro-I<sub>1</sub>-Leu-Pro-J<sub>1</sub>-<sup>30</sup>Thr-K<sub>1</sub>-Val-Gly-Ser-<sup>35</sup>Asn-Thr-Tyr-Z  
wherein

A<sub>1</sub> is Lys, Ala, Ser or Hydrogen;

B<sub>1</sub> is Ala, Set or Thr;

C<sub>1</sub> is Val, Leu or Ile;

D<sub>1</sub> is His or Arg;

E<sub>1</sub> is Ser or Thr;

F<sub>1</sub> is Ser, Thr, Gln or Asn;

G<sub>1</sub> is Asn, Gln or His;

H<sub>1</sub> is Phe, Leu or Tyr;

I<sub>1</sub> is Ile, Val, Ala or Leu;

J<sub>1</sub> is Ser, Pro or Thr;

K<sub>1</sub> is Asn, Asp or Gln;

X and Y are independently selected residues having side chains which are chemically bonded to each other to form an intramolecular linkage, wherein said intramolecular linkage comprises a

disulfide bond, a lactam or a thioether linkage; and Z is an amino, alkylamino, dialkylamino, cycloalkylamino, arylamino, aralkylamino, alkyloxy, aryloxy or aralkyloxy; and provided that when A<sub>1</sub> is Lys, B<sub>1</sub> is Ala, C<sub>1</sub> is Val, D<sub>1</sub> is Arg, E<sub>1</sub> is Ser, F<sub>1</sub> is Ser, G<sub>1</sub> is Asn, H<sub>1</sub> is Leu, I<sub>1</sub> is Val, J<sub>1</sub> is Pro, and K<sub>1</sub> is Asn; then one or more A<sub>1</sub> to K<sub>1</sub> is a D-amino acid and Z is selected from the group consisting of alkylamino, dialkylamino, cycloalkylamino, arylamino, aralkylamino, alkyloxy, aryloxy or aralkyloxy.

26. (Previously presented) The method of claim 24 wherein the amylin agonist is an amylin agonist analogue having the following amino acid sequence:

<sup>1</sup>A<sub>1</sub>-X-Asn-Thr-<sup>5</sup>Ala-Thr-Y-Ala-Thr-<sup>10</sup>Gln-Arg-Leu-B<sub>1</sub>-Asn-<sup>15</sup>Phe-Leu-C<sub>1</sub>-D<sub>1</sub>-E<sub>1</sub>-<sup>20</sup>F<sub>1</sub>-G<sub>1</sub>-Asn-H<sub>1</sub>-Gly-<sup>25</sup>Pro-I<sub>1</sub>-Leu-J<sub>1</sub>-Pro-<sup>30</sup>Thr-K<sub>1</sub>-Val-Gly-Ser-<sup>35</sup>Asn-Thr-Tyr-Z

wherein

A<sub>1</sub> is Lys, Ala, Ser or hydrogen;

B<sub>1</sub> is Ala, Ser or Thr;

C<sub>1</sub> is Val, Leu or Ile;

D<sub>1</sub> is His or Arg;

E<sub>1</sub> is Ser or Thr;

F<sub>1</sub> is Ser, Thr, Gln or Asn;

G<sub>1</sub> is Asn, Gln or His;

H<sub>1</sub> is Phe, Leu or Tyr;

I<sub>1</sub> is Ile, Val, Ala or Leu;

J<sub>1</sub> is Ser, Pro, Leu, Ile or Thr;

K<sub>1</sub> is Asn, Asp or Gln;

X and Y are independently selected residues having side chains which are chemically bonded to each other to form an intramolecular linkage, wherein said intramolecular linkage comprises a disulfide bond, a lactam or a thioether linkage; and Z is amino, alkylamino, dialkylamino, cycloalkylamino, arylamino, aralkylamino, alkyloxy, aryloxy or aralkyloxy, and provided that when

(a) A<sub>1</sub> is Lys, B<sub>1</sub> is Ala, C<sub>1</sub> is Val, D<sub>1</sub> is Arg, E<sub>1</sub> is Ser, F<sub>1</sub> is Ser, G<sub>1</sub> is Asn, H<sub>1</sub> is Leu, I<sub>1</sub> is Val, J<sub>1</sub> is Pro and K<sub>1</sub> is Asn; or

(b) A<sub>1</sub> is Lys, B<sub>1</sub> is Ala, C<sub>1</sub> is Val, D<sub>1</sub> is His, E<sub>1</sub> is Ser, F<sub>1</sub> is Asn, G<sub>1</sub> is Asn, H<sub>1</sub> is Leu, I<sub>1</sub> is Val, J<sub>1</sub> is Ser and K<sub>1</sub> is Asn; then one or more of A<sub>1</sub> to K<sub>1</sub> is a D-amino acid and Z is selected from the group consisting of alkylamino, dialkylamino, cycloalkylamino, arylamino, aralkylamino, alkyloxy, aryloxy or aralkyloxy.

27. (Previously presented) The method of claim 24 wherein the amylin agonist is an amylin agonist analogue having the following amino acid sequence:

<sup>1</sup>A<sub>1</sub>-X-Asn-Thr-<sup>5</sup>Ala-Thr-Y-Ala-Thr-<sup>10</sup>Gln-Arg-Leu-B<sub>1</sub>-Asn-<sup>15</sup>Phe-Leu-C<sub>1</sub>-D<sub>1</sub>-E<sub>1</sub>-<sup>20</sup>F<sub>1</sub>-G<sub>1</sub>-Asn-H<sub>1</sub>-Gly-<sup>25</sup>I<sub>1</sub>-J<sub>1</sub>-Leu-Pro-Pro-<sup>30</sup>Thr-K<sub>1</sub>-Val-Gly-Ser-<sup>35</sup>Asn-Thr-Tyr-Z

wherein

A<sub>1</sub> is Lys, Ala, Ser or hydrogen;

B<sub>1</sub> is Ala, Ser or Thr;

C<sub>1</sub> is Val, Leu or Ile;

D<sub>1</sub> is His or Arg;

E<sub>1</sub> is Ser or Thr;

F<sub>1</sub> is Ser, Thr, Gln or Asn;

G<sub>1</sub> is Asn, Gln or His;

H<sub>1</sub> is Phe, Leu or Tyr;

I<sub>1</sub> is Ala or Pro;

J<sub>1</sub> is Ile, Val, Ala or Leu;

K<sub>1</sub> is Asn, Asp or Gln;

X and Y are independently selected residues having side chains which are chemically bonded to each other to form an intramolecular linkage, wherein said intramolecular linkage comprises a disulfide bond, a lactam or a thioether linkage; and Z is amino, alkylamino, dialkylamino, cycloalkylamino, arylamino, aralkylamino, alkyloxy, aryloxy or aralkyloxy; and provided that when A<sub>1</sub> is Lys, B<sub>1</sub> is Ala, C<sub>1</sub> is Val, D<sub>1</sub> is Arg, E<sub>1</sub> is Ser, F<sub>1</sub> is Ser, G<sub>1</sub> is Asn, H<sub>1</sub> is Leu, I<sub>1</sub> is Pro, J<sub>1</sub> is Val and K<sub>1</sub> is Asn; then one or more of A<sub>1</sub> to K<sub>1</sub> is a D-amino acid and Z is selected from the group consisting of alkylamino, dialkylamino, cycloalkylamino, arylamino, aralkylamino, alkyloxy, aryloxy or aralkyloxy.

28. (Previously presented) The method of claim 24 wherein the amylin agonist is an amylin agonist analogue having the following amino acid sequence:

<sup>1</sup>A<sub>1</sub>-X-Asn-Thr-<sup>5</sup>Ala-'Ihr-Y-Ala-Thr-<sup>10</sup>Gln-Arg-Leu-B<sub>1</sub>-Asn-<sup>15</sup>Phe-Leu-C<sub>1</sub>-D<sub>1</sub>-E<sub>1</sub>-<sup>20</sup>F<sub>1</sub>-G<sub>1</sub>-Asn-H<sub>1</sub>-Gly-<sup>25</sup>Pro-I<sub>1</sub>-Leu-Pro-Pro-<sup>30</sup>Thr-J<sub>1</sub>-Val-Gly-Ser-<sup>35</sup>Asn-Thr-Tyr-Z

wherein

A<sub>1</sub> is Lys, Ala, Ser or hydrogen;

B<sub>1</sub> is Ala, Ser or Thr;

C<sub>1</sub> is Val, Leu or Ile;

D<sub>1</sub> is His or Arg;

E<sub>1</sub> is Ser or Thr;

F<sub>1</sub> is Ser, Thr, Gln or Asn;

G<sub>1</sub> is Asn, Gln or His;

H<sub>1</sub> is Phe, Leu or Tyr;

I<sub>1</sub> is Ile, Val, Ala or Leu

J<sub>1</sub> is Asn, Asp or Gln;

X and Y are independently selected residues having side chains which are chemically bonded to each other to form an intramolecular linkage wherein said intramolecular linkage comprises a disulfide bond, a lactam or a thioether linkage; and Z is amino, alkylamino, dialkylamino, cycloalkylamino, arylamino, aralkylamino, alkyloxy, aryloxy or aralkyloxy; and provided that when A<sub>1</sub> is Lys, B<sub>1</sub> is Ala, C<sub>1</sub> is Val, D<sub>1</sub> is Arg, E<sub>1</sub> is Ser, F<sub>1</sub> is Ser, G<sub>1</sub> is Asn, H<sub>1</sub> is Leu, I<sub>1</sub> is Val and J<sub>1</sub> is Asn; then one or more of A<sub>1</sub> to J<sub>1</sub> is a D-amino acid and Z is selected from the group consisting of alkylamino, dialkylamino, cycloalkylamino, arylamino, aralkylamino, alkyloxy, aryloxy or aralkyloxy.

29. (Previously presented) The method of claim 24 wherein said amylin agonist is any one of <sup>18</sup>Arg<sup>25,28</sup>Pro-h-amylin, des-<sup>1</sup>Lys<sup>18</sup>Arg<sup>25,28</sup>Pro-h-amylin, <sup>25,28,29</sup>Pro-h-amylin, des-<sup>1</sup>Lys<sup>25,28,29</sup>Pro-h-amylin, <sup>18</sup>Arg<sup>25,28,29</sup>Pro-h-amylin, des-<sup>1</sup>Lys<sup>18</sup>Arg<sup>25,28,29</sup>Pro-h-amylin, <sup>25</sup>Pro<sup>26</sup>Val<sup>28,29</sup>Pro-h-amylin, or des-<sup>1</sup>Lys<sup>25</sup>Pro<sup>26</sup>Val<sup>28,29</sup>Pro-h-amylin.

30. (Previously presented) The method of claim 24 wherein the amylin agonist is  
<sup>25,28,29</sup>Pro-h-amylin.

31-37. (Canceled)

38. (Previously presented) The method of claim 24 wherein the mammal has diabetes.

39. (Previously presented) The method of claim 38 wherein the diabetes is type 1.

40. (Previously presented) The method of claim 38 wherein the diabetes is type 2.

41. (Previously presented) The method of claim 25 wherein the mammal has diabetes.

42. (Previously presented) The method of claim 41 wherein the diabetes is type 1.

43. (Previously presented) The method of claim 41 wherein the diabetes is type 2.

44. (Previously presented) The method of claim 26 wherein the mammal has diabetes.

45. (Previously presented) The method of claim 44 wherein the diabetes is type 1.

46. (Previously presented) The method of claim 44 wherein the diabetes is type 2.

47. (Previously presented) The method of claim 27 wherein the mammal has diabetes.

48. (Previously presented) The method of claim 47 wherein the diabetes is type 1.

49. (Previously presented) The method of claim 47 wherein the diabetes is type 2.

50. (Previously presented) The method of claim 28 wherein the mammal has diabetes.

51. (Previously presented) The method of claim 50 wherein the diabetes is type 1.
52. (Previously presented) The method of claim 50 wherein the diabetes is type 2.
53. (Previously presented) The method of claim 30 wherein the mammal has diabetes.
54. (Previously presented) The method of claim 53 wherein the diabetes is type 1.
55. (Previously presented) The method of claim 53 wherein the diabetes is type 2.
56. (Previously presented) The method of claim 24 wherein the amylin agonist is an amylin agonist analogue having the following amino acid sequence:

$^1\text{A}_1\text{-X-Asn-Thr}^5\text{Ala-Thr-X-Ala-Thr}^{10}\text{Gln-Arg-Leu-B}_1\text{-Asn}^{15}\text{Phe-Leu-C}_1\text{-D}_1\text{-E}_1\text{-}^{20}\text{F}_1\text{-G}_1\text{-Asn-H}_1\text{-Gly}^{25}\text{I}_1\text{-J}_1\text{-Leu-K}_1\text{-L}_1\text{-}^{30}\text{Thr-M}_1\text{-Val-Gly-Ser}^{35}\text{Asn-Thr-Tyr-Z}$

wherein

$\text{A}_1$  is Lys, Ala, Ser, Hydrogen or acetylated Lys;

$\text{B}_1$  is Ala, Ser or Thr;

$\text{C}_1$  is Val, Leu or Ile;

$\text{D}_1$  is His or Arg;

$\text{E}_1$  is Ser or Thr;

$\text{F}_1$  is Ser, Thr, Gln or Asn;

$\text{G}_1$  is Asn, Gln or His;

$\text{H}_1$  is Phe, Leu or Tyr,

$\text{I}_1$  is Ala or Pro;

$\text{J}_1$  is Ile, Val, Ala or Leu;

$\text{K}_1$  is Ser, Pro, Leu, Ile or Thr;

$\text{L}_1$  is Ser, Pro or Thr;

$\text{M}_1$  is Asn, Asp or Gln;

X and Y are independently selected residues having side chains which are chemically bonded to

each other to form an intramolecular linkage, wherein said intramolecular linkage comprises a disulfide bond, a lactam or a thioether linkage; and Z is an amino, alkylamino, dialkylamino, cycloalkylamino, arylamino, aralkylamino, alkyloxy, aryloxy or aralkyloxy; and provided that

(a) when A<sub>1</sub> is Lys, B<sub>1</sub> is Ala, C<sub>1</sub> is Val, D<sub>1</sub> is His, E<sub>1</sub> is Ser, F<sub>1</sub> is Ser, G<sub>1</sub> is Asn, H<sub>1</sub> is Phe, I<sub>1</sub> is Ala, J<sub>1</sub> is Ile, K<sub>1</sub> is Ser, L<sub>1</sub> is Ser, and M<sub>1</sub> is Asn;

(b) when A<sub>1</sub> is Lys, B<sub>1</sub> is Ala, C<sub>1</sub> is Ile, D<sub>1</sub> is Arg, E<sub>1</sub> is Ser, F<sub>1</sub> is Ser, G<sub>1</sub> is Asn, H<sub>1</sub> is Leu, I<sub>1</sub> is Ala, J<sub>1</sub> is Ile, K<sub>1</sub> is Ser, L<sub>1</sub> is Pro, and M<sub>1</sub> is Asn;

(c) when A<sub>1</sub> is Lys, B<sub>1</sub> is Ala, C<sub>1</sub> is Val, D<sub>1</sub> is Arg, E<sub>1</sub> is Thr, F<sub>1</sub> is Ser, G<sub>1</sub> is Asn, H<sub>1</sub> is Leu, I<sub>1</sub> is Ala, J<sub>1</sub> is Ile, K<sub>1</sub> is Ser, L<sub>1</sub> is Pro, and M<sub>1</sub> is Asn;

(d) when A<sub>1</sub> is Lys, B<sub>1</sub> is Ala, C<sub>1</sub> is Val, D<sub>1</sub> is Arg, E<sub>1</sub> is Ser, F<sub>1</sub> is Ser, G<sub>1</sub> is Asn, H<sub>1</sub> is Leu, I<sub>1</sub> is Pro, J<sub>1</sub> is Val, K<sub>1</sub> is Pro, L<sub>1</sub> is Pro, and M<sub>1</sub> is Asn;

(e) when A<sub>1</sub> is Lys, B<sub>1</sub> is Ala, C<sub>1</sub> is Val, D<sub>1</sub> is His, E<sub>1</sub> is Ser, F<sub>1</sub> is Asn, G<sub>1</sub> is Asn, H<sub>1</sub> is Leu, I<sub>1</sub> is Pro, J<sub>1</sub> is Val, K<sub>1</sub> is Ser, L<sub>1</sub> is Pro and M<sub>1</sub> is Asn; or

(f) when A<sub>1</sub> is Lys, B<sub>1</sub> is Thr, C<sub>1</sub> is Val, D<sub>1</sub> is Arg, E<sub>1</sub> is Ser, F<sub>1</sub> is Ser, G<sub>1</sub> is His, H<sub>1</sub> is Leu, I<sub>1</sub> is Ala, J<sub>1</sub> is Ala, K<sub>1</sub> is Leu, L<sub>1</sub> is Pro and M<sub>1</sub> is Asp;

then one or more of any of A<sub>1</sub> to M<sub>1</sub> is not an L-amino acid and Z is not amino.

57. (Previously presented) The method of claim 56 wherein the mammal has diabetes.

58. (Previously presented) The method of claim 57 wherein the diabetes is type 1.

59. (Previously presented) The method of claim 57 wherein the diabetes is type 2.

60-69. (Canceled)